

SB-0008.ST25.txt
SEQUENCE LISTING

<110> Finlay, Brett
Gruenheid, Samantha
Deng, Wanyin
Vallance, Bruce
Puente, Jose L.

<120> BACTERIAL VIRULENCE FACTORS AND USES THEREOF

<130> S&B-0008

<150> PCT/CA2004/001891
<151> 2004-10-29

<150> US 60/515,703
<151> 2003-10-31

<160> 84

<170> PatentIn version 3.3

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<213> Citrobacter rodentium

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atatatgata agaagtataat atccggata accagaggag tagctgaact aaaacaggaa	180
gaatttgtt acgagaaagc cagacggttt tcttatatacg agactatgta ttctgtatgt	240
ccagaagcgt ttgaacctat ttccagaaat gaagccagta cacccggagg aagctggcta	300
acagttataat ccggaaaacg cccaatgggg cagtttctg tagatagttt atacaatcct	360
gatttacatg cattatgtga gcttccggac atttgtgtc agatcttccc taaagaaaat	420
aatgattttt tatacatatgt tgggtgtac agaaatgaca gccctctagg agaacaacgg	480
gcaaataatgt ttatagaatt atataatataaaaagagata tcgtgcaggaa attaaattat	540
gagttaccag agttaaaggc agttaaatct gaaatgatta tcgcacgtga aatgggagaa	600
atcttagct acatgcctgg ggaaatagac agttatatacg aatacataaaa taataaactt	660
tctaaaattt agtag	675

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<210> 17
 <211> 675
 <212> DNA
 <213> Enterohemorrhagic E. coli

<400> 17	
atgattaatc ctgttactaa tactcagggc gtgtccctta taaaatactaa atatgctgaa	60
catgtggtaaaaatattt cccggaaattt aaacatgatt actttatga atcaccat	120
atatatgata agaagtataat atccggata accagaggag tagctgaact aaaacaggaa	180
gaatttgtta acgagaaagc cagacggttt tcttatatga agactatgtt ttctgtatgt	240
ccagaagcgt ttgaacctat ttccagaaat gaagccagta caccggagg aagctggcta	300
acagttatcccgaaaaacg cccaatgggg cagtttctg tagatagttt atacaatcct	360
gatttacatg cattatgtga gcttccggac atttgttgta agatcttccc taaagaaaaat	420
aatgattttt tatacatagt ttttgttac agaaatgaca gccctctagg agaacaacgg	480
gcaaataatgttataatataaaaagagata tcatgcagga attaaattat	540
gagttaccag agttaaaggc agttaaatct gaaatgatta tcgcacgtga aatggagaa	600
atcttagct acatgcctgg ggaaatagac agttatatga aatacataaaa taataaactt	660
tctaaaattt agtag	675

<210> 18
 <211> 570
 <212> DNA
 <213> Citrobacter rodentium

<400> 18	
atgttaccaa caagtggttc ttccagcaaat ctttactcat ggatgtatcttcaggaaaa	60
gagaatccctt cgactccgga atcagtaatg gaacttaatc ataatcattt tctttctcct	120
gaatttacagg agaaactgga ttgttatgttc gccatatattt catgtgccag aaacaatgtat	180
gagcgtgaga atatttaccc ggagcttaagg gatttgttaa gtagcctaattt ggataagaga	240
aacaatgtgtt tgaggtgtat aaatgaagat actgatgagg tgaccggagc tctgagagcg	300
ggaatgacga tagaggacag ggatagttt atcagggatc tttttttctt gcattcattt	360
aaagtaaaaa ttgagggaaatg cagacaagat aaagaggatt ggaaatgttaa agtttatgtat	420
ctgctatgtc cgcatcattt ttcagagcta tatggggatc tacggcaat caaatgcctc	480
gttgaaggat gcagtgtatga ttttagtcctt tttgataacta ttaaggtgcc ggatcttact	540
tacaacaaatgatctttaca atgtggatga	570

<210> 19
 <211> 519
 <212> DNA
 <213> Citrobacter rodentium

<400> 19	
agcaaatctt tactcatgga tgttatatctc aggaaaagag aatccttcga ctccggaaatc	60
agtaagtgaa cttaatcata atcattttctt ttctcctgaa ttacaggaga aactggatgt	120

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tatgttcgcc	atatattcat	gtgccagaaa	caatgatgag	cgtgagaata	tttacccgga	180
gctaaggat	ttttaagta	gcctaatgga	taagagaaac	aatgtgttg	aggtgataaa	240
tgaagatact	gatgaggtga	ccggagctct	gagagcggga	atgacgata	aggacaggg	300
tagttatatc	agggatctt	ttttctgca	ttcattgaaa	gtaaaaattg	aggaaagcag	360
acaagataaa	gaggattgga	aatgtaaagt	ttatgatctg	ctatgtccgc	atcattctc	420
agagctatat	ggggatctac	gggcaatcaa	atgcctcg	tttgcgtt	gaaggatgca	480
tagtccttt	gatactatta	aggtgccgga	tcttactta			519

<210> 20
<211> 570
<212> DNA
<213> Enteropathogenic E. coli

<400>	20						
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ggtaacc	ttt	cgactcc	atc	agtaagt	gagcttaatc	ataatcactt	120
gaattacaag	at	aaacttga	tgttatgg	tc	tatattatt	catgtgccag	180
gagcttgagg	aa	attttca	agagctaagt	gctttgtaa	gtgggctgat	ggataagaga	240
aatagtgtat	tt	gaggtgag	aaatgaaaat	actgatgagg	ttgtcgg	gagc gctgagg	300
ggaatgacga	ta	gaggatag	ggatagttat	atcagggatc	tttttttct	gcattcattg	360
aaagtaaaaa	tt	gagaaag	tagacaaggc	aaagaagatt	cgaatgtaa	agtttataat	420
ctgctatgtc	cg	catcactc	ttcagagcta	tatgg	tgatc	tacgagcaat	480
gtggaaggat	gc	agtgatga	tttaatcct	tttgatatta	ttagggtacc	agatcttact	540
tacaacaaag	gat	ctttaca	atgtggatga				570

<210> 21
<211> 570
<212> DNA
<213> Enterohemorrhagic E. coli

<400>	21						
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ggtaacc	ttt	cgactcc	atc	agtaagt	gagcttaatc	ataatcactt	120
gaattacaag	at	aaacttga	tgttatgg	tc	tatattatt	catgtgccag	180
gagcttgagg	aa	attttca	agagctaagt	gctttgtaa	gtgggctgat	ggataagaga	240
aatagtgtat	tt	gaggtgag	aaatgaaaat	actgatgagg	ttgtcgg	gagc gctgagg	300
ggaatgacga	ta	gaggacag	ggatagttat	atcagggatc	tttttttct	gcattcattg	360
aaagtaaaaa	tt	gagaaag	tagacaaggc	aaagaagatt	cgaatgtaa	agtttataat	420
ctgctatgtc	cg	catcactc	ttcagagcta	tatgg	tgatc	tacgagcaat	480
gtggaaggat	gc	agtgatga	tttaatcct	tttgatatta	ttagggtacc	agatcttact	540
tacaacaaag	gat	ctttaca	atgtggatga				570

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<210> 22
<211> 430
<212> PRT
<213> Citrobacter rodentium

<400> 22

Met Asn Ile Gln Pro Asn Ile His Ser Gly Ile Thr Thr Gln Asn Asn
1 5 10 15

Gln Gln His His His Ala Glu Gln Val Pro Val Ser Ser Ser Ile Pro
20 25 30

Arg Ser Asp Leu Pro Pro Asn Cys Glu Ala Gly Phe Val Val His Ile
35 40 45

Pro Glu Asp Ile Gln Gln His Val Pro Glu Cys Gly Glu Thr Thr Ala
50 55 60

Leu Leu Ser Leu Ile Lys Asp Glu Gly Leu Leu Ser Gly Leu Asp Lys
65 70 75 80

Tyr Leu Ala Pro His Leu Glu Glu Gly Ser Leu Gly Lys Lys Ala Leu
85 90 95

Asp Thr Phe Gly Leu Phe Asn Val Thr Gln Met Ala Leu Glu Ile Pro
100 105 110

Ser Ser Val Pro Gly Ile Ser Gly Lys Tyr Gly Val Gln Met Asn Ile
115 120 125

Val Lys Pro Asp Ile His Pro Thr Thr Gly Asn Tyr Phe Leu Gln Leu
130 135 140

Phe Pro Leu His Asp Glu Ile Gly Phe Asn Phe Lys Asp Leu Pro Gly
145 150 155 160

Pro Leu Lys Asn Ala Leu Thr Asn Ser Ser Ile Ser Ala Thr Ala Ser
165 170 175

Thr Val Ala Pro Thr Pro Asn Asp Pro Met Pro Trp Phe Gly Leu Thr
180 185 190

Ala Gln Val Val Arg Asn His Gly Val Glu Leu Pro Ile Val Lys Thr
195 200 205

Glu Asn Gly Trp Lys Leu Val Gly Glu Thr Pro Leu Thr Pro Asp Gly
210 215 220

Pro Lys Ala Asn Tyr Thr Glu Glu Trp Val Ile Arg Pro Gly Glu Ala
225 230 235 240

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Asp Phe Lys Tyr Gly Thr Ser Pro Leu Gln Ala Thr Leu Gly Leu Glu
245 250 255

Phe Gly Ala His Phe Lys Trp Asp Leu Asp Asn Pro Asn Thr Lys Tyr
260 265 270

Ala Ile Leu Thr Asn Ala Ala Asn Ala Ile Gly Ala Ala Gly Gly
275 280 285

Phe Ala Val Ser Lys Val Pro Gly Ile Asp Pro Met Leu Ser Pro His
290 295 300

Val Gly Ala Met Leu Gly Gln Ala Ala Gly His Ala Val Gln Cys Asn
305 310 315 320

Thr Pro Gly Leu Lys Pro Asp Thr Ile Leu Trp Trp Ala Gly Ala Thr
325 330 335

Phe Gly Ala Ala Asp Leu Asn Lys Ala Glu Phe Asp Lys Val Arg Phe
340 345 350

Thr Asp Tyr Pro Arg Ile Trp Phe His Ala Arg Glu Gly Ala Leu Phe
355 360 365

Pro Asn Lys Gln Asp Ile Ala Arg Val Thr Gly Ala Asp Ile Lys Ala
370 375 380

Met Glu Glu Gly Val Pro Val Gly His Gln His Pro Lys Pro Glu Asp
385 390 395 400

Val Val Ile Asp Ile Glu Gly Asn Ser Pro His His Asn Pro Ser
405 410 415

Asn Tyr Val Asp Thr Phe Glu Ile Ile Gln Glu Thr Arg Val
420 425 430

<210> 23
<211> 440
<212> PRT
<213> Enteropathogenic E. coli

<400> 23

Met Asn Ile Gln Pro Ile Val Thr Ser Gly Ile Thr Thr Gln Asn Asn
1 5 10 15

Arg His His His Ala Glu Gln Thr Ser Pro Thr Gln Ile Pro Gln Ser
20 25 30

Glu Leu Pro Asn Gly Cys Glu Thr Gly Phe Val Val His Ile Pro Glu
35 40 45

Asp Met Gln Arg His Ala Pro Glu Cys Gly Glu Thr Thr Ala Leu Leu
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50

55

60

Ser Leu Ile Lys Asp Glu Gly Leu Leu Ser Gly Leu Asp Lys Tyr Leu
65 70 75 80

Ala Pro His Leu Glu Glu Gly Ser Ala Gly Lys Lys Ala Leu Asp Met
85 90 95

Phe Gly Leu Phe Asn Val Ser Gln Met Ala Leu Glu Ile Pro Ser Thr
100 105 110

Val Pro Gly Ile Ser Gly Lys Tyr Gly Val Gln Leu Asn Ile Val Lys
115 120 125

Pro Asp Ile His Pro Thr Ser Gly Asn Tyr Phe Leu Gln Ile Phe Pro
130 135 140

Leu His Asp Glu Ile Gly Ile Asn Phe Lys Asp Leu Pro Gly Pro Leu
145 150 155 160

Lys Asn Ala Leu Ser Asn Ser Asn Ile Pro Thr Thr Val Ser Thr Ala
165 170 175

Ala Ser Thr Ile Ala Ser Ala Thr Thr Ser Thr Val Thr Thr Ala Ser
180 185 190

Lys Asp Pro Ile Pro Trp Phe Gly Leu Thr Ala Gln Val Val Arg Asn
195 200 205

His Gly Val Glu Leu Pro Ile Val Lys Thr Glu Asn Gly Trp Lys Leu
210 215 220

Val Gly Glu Thr Pro Leu Thr Pro Asp Gly Pro Lys Ala Asn Tyr Thr
225 230 235 240

Glu Glu Trp Val Ile Arg Pro Gly Glu Ala Asp Phe Lys Tyr Gly Ala
245 250 255

Ser Pro Leu Gln Ala Thr Leu Gly Leu Glu Phe Gly Ala His Phe Lys
260 265 270

Trp Asp Leu Asp Asn Pro Asn Thr Lys Tyr Ala Val Leu Thr Asn Ala
275 280 285

Ala Ala Asn Ala Leu Gly Ala Val Gly Gly Phe Ala Val Ser Arg Phe
290 295 300

Thr Gly Thr Asp Pro Met Leu Ser Pro His Ile Gly Ala Met Val Gly
305 310 315 320

Gln Ala Ala Gly His Ala Ile Gln Tyr Asn Thr Pro Gly Leu Lys Pro
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335

Asp Thr Ile Leu Trp Trp Ala Gly Thr Thr Leu Gly Leu Ala Asp Leu
 340 345 350

Asn Lys Ala Glu Phe Gly Glu Ala Arg Phe Thr Asp Tyr Pro Arg Ile
 355 360 365

Trp Trp His Ala Arg Glu Gly Ala Ile Phe Pro Asn Lys Ala Asp Ile
 370 375 380

Glu His Ala Thr Gly Ala Asp Ile Arg Ala Met Glu Glu Gly Val Ser
 385 390 395 400

Val Gly Gln Arg His Pro Asn Pro Glu Asp Val Val Ile Asn Ile Glu
 405 410 415

Ser Asn Asn Ser Pro His His Asn Pro Ser Asn Tyr Val Asp Thr Val
 420 425 430

Asp Ile Ile Gln Glu Thr Arg Val
 435 440

<210> 24

<211> 441

<212> PRT

<213> Enterohemorrhagic E. coli

<400> 24

Met Asn Ile Gln Pro Thr Ile Gln Ser Gly Ile Thr Ser Gln Asn Asn
 1 5 10 15

Gln His His Gln Thr Glu Gln Ile Pro Ser Thr Gln Ile Pro Gln Ser
 20 25 30

Glu Leu Pro Leu Gly Cys Gln Ala Gly Phe Val Val Asn Ile Pro Asp
 35 40 45

Asp Ile Gln Gln His Ala Pro Glu Cys Gly Glu Thr Thr Ala Leu Leu
 50 55 60

Ser Leu Ile Lys Asp Lys Gly Leu Leu Ser Gly Leu Asp Glu Tyr Ile
 65 70 75 80

Ala Pro His Leu Glu Glu Gly Ser Ile Gly Lys Lys Thr Leu Asp Met
 85 90 95

Phe Gly Leu Phe Asn Val Thr Gln Met Ala Leu Glu Ile Pro Ser Ser
 100 105 110

Val Ser Gly Ile Ser Gly Lys Tyr Gly Val Gln Leu Asn Ile Val Lys
 115 120 125

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Pro Asp Ile His Pro Thr Ser Gly Asn Tyr Phe Leu Gln Ile Phe Pro
130 135 140

Leu His Asp Glu Ile Gly Phe Asn Phe Lys Asp Leu Pro Gly Pro Leu
145 150 155 160

Lys Asn Ala Leu Ser Asn Ser Asn Ile Ser Thr Thr Ala Val Ser Thr
165 170 175

Ile Ala Ser Thr Gly Thr Ser Ala Thr Thr Ser Thr Val Thr Thr Glu
180 185 190

Pro Lys Asp Pro Ile Pro Trp Phe Gly Leu Thr Ala Gln Val Val Arg
195 200 205

Asn His Gly Val Glu Leu Pro Ile Val Lys Thr Glu Asn Gly Trp Lys
210 215 220

Leu Val Gly Glu Thr Pro Leu Thr Pro Asp Gly Pro Lys Ala Asn Tyr
225 230 235 240

Thr Glu Glu Trp Val Ile Arg Pro Gly Glu Ala Asp Phe Lys Tyr Gly
245 250 255

Ala Ser Pro Leu Gln Ala Thr Leu Gly Leu Glu Phe Gly Ala His Phe
260 265 270

Lys Trp Asp Leu Asp Asn Pro Asn Thr Lys Tyr Ala Val Leu Thr Asn
275 280 285

Ala Ala Ala Asn Ala Leu Gly Ala Leu Gly Gly Phe Ala Val Ser Arg
290 295 300

Phe Ala Ser Thr Asp Pro Met Leu Ser Pro His Ile Gly Ala Met Val
305 310 315 320

Gly Gln Ala Ala Gly His Ala Ile Gln Tyr Asn Thr Pro Gly Leu Lys
325 330 335

Pro Asp Thr Ile Leu Trp Trp Ala Gly Ala Thr Leu Gly Ala Ala Asp
340 345 350

Leu Asn Lys Ala Glu Phe Glu Val Ala Arg Phe Thr Asp Tyr Pro Arg
355 360 365

Ile Trp Trp His Ala Arg Glu Gly Ala Ile Phe Pro Asn Lys Ala Asp
370 375 380

Ile Glu His Ala Thr Gly Ala Asp Ile Arg Ala Met Glu Glu Gly Ile
385 390 395 400

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Pro Val Gly Gln Arg His Pro Asn Pro Glu Asp Val Val Ile Asp Ile
405 410 415

Glu Ser Asn Gly Leu Pro His His Asn Pro Ser Asn His Val Asp Ile
420 425 430

Phe Asp Ile Ile Gln Glu Thr Arg Val
435 440

<210> 25
<211> 204
<212> PRT
<213> Citrobacter rodentium

<400> 25

Ile Leu Phe Gln Trp Phe Glu Ala Arg Pro Glu Arg Tyr Gly Lys Gly
1 5 10 15

Glu Val Pro Ile Leu Asn Thr Lys Glu His Pro Tyr Leu Ser Asn Ile
20 25 30

Ile Asn Ala Ala Lys Ile Glu Asn Glu Arg Val Ile Gly Val Leu Val
35 40 45

Asp Gly Asp Phe Thr Tyr Glu Gln Arg Lys Glu Phe Leu Ser Leu Glu
50 55 60

Asp Glu His Gln Asn Ile Lys Ile Tyr Arg Glu Asn Val Asp Phe
65 70 75 80

Ser Met Tyr Asp Lys Lys Leu Ser Asp Ile Tyr Leu Glu Asn Ile His
85 90 95

Glu Gln Glu Ser Tyr Pro Ala Ser Glu Arg Asp Asn Tyr Leu Leu Gly
100 105 110

Leu Leu Arg Glu Glu Leu Lys Asn Ile Pro Tyr Gly Lys Asp Ser Leu
115 120 125

Ile Glu Ser Tyr Ala Glu Lys Arg Gly His Thr Trp Phe Asp Phe Phe
130 135 140

Arg Asn Leu Ala Val Leu Lys Gly Gly Leu Phe Thr Glu Thr Gly
145 150 155 160

Lys Thr Gly Cys His Asn Ile Ser Pro Cys Gly Gly Cys Ile Tyr Leu
165 170 175

Asp Ala Asp Met Ile Ile Thr Asp Lys Leu Gly Val Leu Tyr Ala Pro
180 185 190

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Asp Gly Ile Ala Val His Val Asp Cys Asn Asp Glu
195 200

<210> 26
<211> 186
<212> PRT
<213> Citrobacter rodentium

<400> 26

Arg Pro Glu Arg Tyr Gly Lys Gly Glu Val Pro Ile Leu Asn Thr Lys
1 5 10 15

Glu His Pro Tyr Leu Ser Asn Ile Ile Asn Ala Ala Lys Ile Glu Asn
20 25 30

Glu Arg Val Ile Gly Val Leu Val Asp Gly Asp Phe Thr Tyr Glu Gln
35 40 45

Arg Lys Glu Phe Leu Ser Leu Glu Asp Glu His Gln Asn Ile Lys Ile
50 55 60

Ile Tyr Arg Glu Asn Val Asp Phe Ser Met Tyr Asp Lys Lys Leu Ser
65 70 75 80

Asp Ile Tyr Leu Glu Asn Ile His Glu Gln Glu Ser Tyr Pro Ala Ser
85 90 95

Glu Arg Asp Asn Tyr Leu Leu Gly Leu Leu Arg Glu Glu Leu Lys Asn
100 105 110

Ile Pro Tyr Gly Lys Asp Ser Leu Ile Glu Ser Tyr Ala Glu Lys Arg
115 120 125

Gly His Thr Trp Phe Asp Phe Phe Arg Asn Leu Ala Val Leu Lys Gly
130 135 140

Gly Gly Leu Phe Thr Glu Thr Gly Lys Thr Gly Cys His Asn Ile Ser
145 150 155 160

Pro Cys Gly Cys Ile Tyr Leu Asp Ala Asp Met Ile Ile Thr Asp
165 170 175

Lys Leu Gly Val Leu Tyr Ala Pro Asp Gly
180 185

<210> 27
<211> 329
<212> PRT
<213> Enteropathogenic E. coli

<400> 27

Met Leu Ser Ser Leu Asn Val Leu Gln Ser Ser Phe Arg Gly Lys Thr
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Ala Leu Ser Asn Ser Thr Leu Leu Gln Lys Val Ser Phe Ala Gly Lys
20 25 30

Glu Tyr Ser Leu Glu Pro Ile Asp Glu Arg Thr Pro Ile Leu Phe Gln
35 40 45

Trp Phe Glu Ala Arg Pro Glu Arg Tyr Glu Lys Gly Glu Val Pro Ile
50 55 60

Leu Asn Thr Lys Glu His Pro Tyr Leu Ser Asn Ile Ile Asn Ala Ala
65 70 75 80

Lys Ile Glu Asn Glu Arg Ile Ile Gly Val Leu Val Asp Gly Asn Phe
85 90 95

Thr Tyr Glu Gln Lys Lys Glu Phe Leu Asn Leu Glu Asn Glu His Gln
100 105 110

Asn Ile Lys Ile Ile Tyr Arg Ala Asp Val Asp Phe Ser Met Tyr Asp
115 120 125

Lys Lys Leu Ser Asp Ile Tyr Leu Glu Asn Ile His Lys Gln Glu Ser
130 135 140

Tyr Pro Ala Ser Glu Arg Asp Asn Tyr Leu Leu Gly Leu Leu Arg Glu
145 150 155 160

Glu Leu Lys Asn Ile Pro Glu Gly Lys Asp Ser Leu Ile Glu Ser Tyr
165 170 175

Ala Glu Lys Arg Glu His Thr Trp Phe Asp Phe Phe Arg Asn Leu Ala
180 185 190

Ile Leu Lys Ala Gly Ser Leu Phe Thr Glu Thr Gly Lys Thr Gly Cys
195 200 205

His Asn Ile Ser Pro Cys Ser Gly Cys Ile Tyr Leu Asp Ala Asp Met
210 215 220

Ile Ile Thr Asp Lys Leu Gly Val Leu Tyr Ala Pro Asp Gly Ile Ala
225 230 235 240

Val His Val Asp Cys Asn Asp Glu Ile Lys Ser Leu Glu Asn Gly Ala
245 250 255

Ile Val Val Asn Arg Ser Asn His Pro Ala Leu Leu Ala Gly Leu Asp
260 265 270

Ile Met Lys Ser Lys Val Asp Ala His Pro Tyr Tyr Asp Gly Leu Gly
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280

285

Lys Gly Ile Lys Arg His Phe Asn Tyr Ser Ser Leu His Asn Tyr Asn
 290 295 300

Ala Phe Cys Asp Phe Ile Glu Phe Lys His Glu Asn Ile Ile Pro Asn
 305 310 315 320

Thr Ser Met Tyr Thr Ser Ser Ser Trp
 325

<210> 28

<211> 329

<212> PRT

<213> Enterohemorrhagic E. coli

<400> 28

Met Leu Ser Ser Leu Asn Val Leu Gln Ser Ser Phe Arg Gly Lys Thr
 1 5 10 15

Ala Leu Ser Asn Ser Thr Leu Leu Gln Lys Val Ser Phe Ala Gly Lys
 20 25 30

Glu Tyr Pro Leu Glu Pro Ile Asp Glu Lys Thr Pro Ile Leu Phe Gln
 35 40 45

Trp Phe Glu Ala Arg Pro Glu Arg Tyr Glu Lys Gly Glu Val Pro Ile
 50 55 60

Leu Asn Thr Lys Glu His Pro Tyr Leu Ser Asn Ile Ile Asn Ala Ala
 65 70 75 80

Lys Ile Glu Asn Glu Arg Ile Ile Gly Val Leu Val Asp Gly Asn Phe
 85 90 95

Thr Tyr Glu Gln Lys Lys Glu Phe Leu Ser Leu Glu Asn Glu Tyr Gln
 100 105 110

Asn Ile Lys Ile Ile Tyr Arg Ala Asp Val Asp Phe Ser Met Tyr Asp
 115 120 125

Lys Lys Leu Ser Asp Ile Tyr Leu Glu Asn Ile His Lys Gln Glu Ser
 130 135 140

Tyr Pro Ala Ser Glu Arg Asp Asn Tyr Leu Leu Gly Leu Leu Arg Glu
 145 150 155 160

Glu Leu Lys Asn Ile Pro Glu Gly Lys Asp Ser Leu Ile Glu Ser Tyr
 165 170 175

Ala Glu Lys Arg Glu His Thr Trp Phe Asp Phe Phe Arg Asn Leu Ala
 180 185 190

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Met Leu Lys Ala Gly Ser Leu Phe Thr Glu Thr Gly Lys Thr Gly Cys
195 200 205

His Asn Ile Ser Pro Cys Ser Gly Cys Ile Tyr Leu Asp Ala Asp Met
210 215 220

Ile Ile Thr Asp Lys Leu Gly Val Leu Tyr Ala Pro Asp Gly Ile Ala
225 230 235 240

Val His Val Asp Cys Asn Asp Glu Ile Lys Ser Leu Glu Asn Gly Ala
245 250 255

Ile Val Val Asn Arg Ser Asn His Pro Ala Leu Leu Ala Gly Leu Asp
260 265 270

Ile Met Lys Ser Lys Val Asp Ala His Pro Tyr Tyr Asp Gly Leu Gly
275 280 285

Lys Gly Ile Lys Arg His Phe Asn Tyr Ser Ser Leu His Asp Tyr Asn
290 295 300

Ala Phe Cys Asp Phe Ile Glu Phe Lys His Glu Asn Ile Ile Pro Asn
305 310 315 320

Thr Ser Met Tyr Thr Cys Ser Ser Trp
325

<210> 29

<211> 326

<212> PRT

<213> Enterohemorrhagic E. coli

<400> 29

Met Leu Ser Pro Ile Arg Thr Thr Phe His Asn Ser Val Asn Ile Val
1 5 10 15

Gln Ser Ser Pro Cys Gln Thr Val Ser Phe Ala Gly Lys Glu Tyr Glu
20 25 30

Leu Lys Val Ile Asp Glu Lys Thr Pro Ile Leu Phe Gln Trp Phe Glu
35 40 45

Pro Asn Pro Glu Arg Tyr Lys Lys Asp Glu Val Pro Ile Val Asn Thr
50 55 60

Lys Gln His Pro Tyr Leu Asp Asn Val Thr Asn Ala Ala Arg Ile Glu
65 70 75 80

Ser Asp Arg Met Ile Gly Ile Phe Val Asp Gly Asp Phe Ser Val Asn
85 90 95

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Gln Lys Thr Ala Phe Ser Lys Leu Glu Arg Asp Phe Glu Asn Val Met
100 105 110

Ile Ile Tyr Arg Glu Asp Val Asp Phe Ser Met Tyr Asp Arg Lys Leu
115 120 125

Ser Asp Ile Tyr His Asp Ile Ile Cys Glu Gln Arg Leu Arg Thr Glu
130 135 140

Asp Lys Arg Asp Glu Tyr Leu Leu Asn Leu Leu Glu Lys Glu Leu Arg
145 150 155 160

Glu Ile Ser Lys Ala Gln Asp Ser Leu Ile Ser Met Tyr Ala Lys Lys
165 170 175

Arg Asn His Ala Trp Phe Asp Phe Phe Arg Asn Leu Ala Leu Leu Lys
180 185 190

Ala Gly Glu Ile Phe Arg Cys Thr Tyr Asn Thr Lys Asn His Gly Ile
195 200 205

Ser Phe Gly Glu Gly Cys Ile Tyr Leu Asp Met Asp Met Ile Leu Thr
210 215 220

Gly Lys Leu Gly Thr Ile Tyr Ala Pro Asp Gly Ile Ser Met His Val
225 230 235 240

Asp Arg Arg Asn Asp Ser Val Asn Ile Glu Asn Ser Ala Ile Ile val
245 250 255

Asn Arg Ser Asn His Pro Ala Leu Leu Glu Gly Leu Ser Phe Met His
260 265 270

Ser Lys Val Asp Ala His Pro Tyr Tyr Asp Gly Leu Gly Lys Gly Val
275 280 285

Lys Lys Tyr Phe Asn Phe Thr Pro Leu His Asn Tyr Asn His Phe Cys
290 295 300

Asp Phe Ile Glu Phe Asn His Pro Asn Ile Ile Met Asn Thr Ser Gln
305 310 315 320

Tyr Thr Cys Ser Ser Trp
325

<210> 30

<211> 330

<212> PRT

<213> Citrobacter rodentium

<400> 30

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Met Lys Ile Pro Ser Leu Gln Pro Ser Phe Asn Phe Phe Ala Pro Ala
 1 5 10 15

Gly Tyr Ser Ala Ala Val Ala Pro Asn Arg Ser Asp Asn Ala Tyr Ala
 20 25 30

Asp Tyr Val Leu Asp Ile Gly Lys Arg Ile Pro Leu Ser Ala Glu Asp
 35 40 45

Leu Gly Asn Leu Tyr Glu Asn Val Ile Arg Ala Val Arg Asp Ser Arg
 50 55 60

Ser Lys Leu Ile Asp Gln His Thr Val Asp Met Ile Gly Asn Thr Ile
 65 70 75 80

Leu Asp Ala Leu Ser Arg Ser Gln Thr Phe Arg Asp Ala Val Ser Tyr
 85 90 95

Gly Ile His Asn Lys Glu Val His Ile Gly Cys Ile Lys Tyr Arg Asn
 100 105 110

Glu Tyr Glu Leu Asn Gly Glu Ser Pro Val Lys Val Asp Asp Ile Gln
 115 120 125

Ser Leu Thr Cys Thr Glu Leu Tyr Glu Tyr Asp Val Gly Gln Glu Pro
 130 135 140

Ile Leu Pro Ile Cys Glu Ala Gly Glu Asn Asp Asn Glu Glu Pro Tyr
 145 150 155 160

Val Ser Phe Ser Val Ala Pro Asp Thr Asp Ser Tyr Glu Met Pro Ser
 165 170 175

Trp Gln Glu Gly Leu Ile His Glu Ile Ile His His Val Thr Gly Ala
 180 185 190

Ser Asp Pro Ser Gly Asp Ser Asn Ile Glu Leu Gly Pro Thr Glu Ile
 195 200 205

Leu Ala Arg Arg Val Ala Gln Glu Leu Gly Trp Thr Val Pro Asp Phe
 210 215 220

Ile Gly Tyr Ala Glu Pro Asp Arg Glu Ala His Leu Arg Gly Arg Asn
 225 230 235 240

Leu Asn Ala Leu Arg Gln Ala Ala Met Arg His Glu Asp Asn Glu Arg
 245 250 255

Thr Phe Phe Glu Arg Leu Gly Met Ile Ser Asp Arg Tyr Glu Ala Ser
 260 265 270

Pro Asp Phe Thr Glu Tyr Ser Ala Val Ser Asn Ile Glu Tyr Gly Phe
 275 280 285

Ile Gln Gln His Asp Phe Pro Gly Leu Ala Ile Asp Asp Asn Leu Gln
 290 295 300

Asp Ala Asn Gln Ile Gln Leu Tyr His Gly Ala Pro Tyr Ile Phe Thr
 305 310 315 320

Phe Gly Asp Val Asp Lys His Asn Gln Arg
 325 330

<210> 31

<211> 330

<212> PRT

<213> Enteropathogenic E. coli

<400> 31

Met Lys Ile Pro Ser Leu Gln Ser Asn Phe Asn Phe Ser Ala Pro Ala
 1 5 10 15

Gly Tyr Ser Ala Pro Ile Ala Pro Asn Arg Ala Glu Asn Ala Tyr Ala
 20 25 30

Asp Tyr Val Leu Asp Ile Gly Lys Arg Ile Pro Leu Ser Ala Ala Asp
 35 40 45

Leu Ser Asn Val Tyr Glu Ser Val Ile Arg Ala Val His Asp Ser Arg
 50 55 60

Ser Arg Leu Ile Asp Gln His Thr Val Asp Met Ile Gly Asn Thr Val
 65 70 75 80

Leu Asp Ala Leu Ser Arg Ser Gln Thr Phe Arg Asp Ala Val Ser Tyr
 85 90 95

Gly Ile His Asn Glu Lys Val His Ile Gly Cys Ile Lys Tyr Arg Asn
 100 105 110

Glu Tyr Glu Leu Asn Glu Glu Ser Ser Val Lys Ile Asp Asp Ile Gln
 115 120 125

Ser Leu Thr Cys Asn Glu Leu Tyr Glu Tyr Asp Val Gly Gln Glu Pro
 130 135 140

Ile Phe Pro Ile Cys Glu Ala Gly Glu Asn Asp Asn Glu Glu Pro Tyr
 145 150 155 160

Val Ser Phe Ser Val Ala Pro Asp Thr Asp Ser Tyr Glu Met Pro Ser
 165 170 175

Trp Gln Glu Gly Leu Ile His Glu Ile Ile His His Val Thr Gly Ser
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180

185

190

Ser Asp Pro Ser Gly Asp Ser Asn Ile Glu Leu Gly Pro Thr Glu Ile
 195 200 205

Leu Ala Arg Arg Val Ala Gln Glu Leu Gly Trp Ser Val Pro Asp Phe
 210 215 220

Lys Gly Tyr Ala Glu Pro Glu Arg Glu Ala His Leu Arg Leu Arg Asn
 225 230 235 240

Leu Asn Ala Leu Arg Gln Ala Ala Met Arg His Glu Glu Asn Glu Arg
 245 250 255

Ala Phe Phe Glu Arg Leu Gly Thr Ile Ser Asp Arg Tyr Glu Ala Ser
 260 265 270

Pro Asp Phe Thr Glu Tyr Ser Ala Val Ser Asn Ile Gly Tyr Gly Phe
 275 280 285

Ile Gln Gln His Asp Phe Pro Gly Leu Ala Ile Asn Asp Asn Leu Gln
 290 295 300

Asp Ala Asn Gln Ile Gln Leu Tyr His Gly Ala Pro Tyr Ile Phe Thr
 305 310 315 320

Phe Gly Asp Val Asp Lys His Asn Gln Arg
 325 330

<210> 32

<211> 330

<212> PRT

<213> Enterohemorrhagic E. coli

<400> 32

Met Lys Ile Pro Ser Leu Gln Ser Asn Phe Asn Phe Ser Ala Pro Ala
 1 5 10 15

Gly Tyr Ser Ala Pro Ile Ala Pro Asn Arg Ala Glu Asn Ala Tyr Ala
 20 25 30

Asp Tyr Val Leu Asp Ile Gly Lys Arg Ile Pro Leu Ser Ala Ala Asp
 35 40 45

Leu Ser Asn Val Tyr Glu Ser Val Ile Arg Ala Val His Asp Ser Arg
 50 55 60

Ser Arg Leu Ile Asp Gln His Thr Val Asp Met Ile Gly Asn Thr Val
 65 70 75 80

Leu Asp Ala Leu Ser Arg Ser Gln Thr Phe Arg Asp Ala Val Ser Tyr
 85 90 95

SB-0008.ST25.txt

Gly Ile His Asn Glu Lys Val His Ile Gly Cys Ile Lys Tyr Arg Asn
100 105 110

Glu Tyr Glu Leu Asn Glu Glu Ser Ser Val Lys Ile Asp Asp Ile Gln
115 120 125

Ser Leu Thr Cys Asn Glu Leu Tyr Glu Tyr Asp Val Gly Gln Glu Pro
130 135 140

Ile Phe Pro Ile Cys Glu Ala Gly Glu Asn Asp Asn Glu Glu Pro Tyr
145 150 155 160

Val Ser Phe Ser Val Ala Pro Asp Thr Asp Ser Tyr Glu Met Pro Ser
165 170 175

Trp Gln Glu Gly Leu Ile His Glu Ile Ile His His Val Thr Gly Ser
180 185 190

Ser Asp Pro Ser Gly Asp Ser Asn Ile Glu Leu Gly Pro Thr Glu Ile
195 200 205

Leu Ala Arg Arg Val Ala Gln Glu Leu Gly Trp Ser Val Pro Asp Phe
210 215 220

Lys Gly Tyr Ala Glu Pro Glu Arg Glu Ala His Leu Arg Leu Arg Asn
225 230 235 240

Leu Asn Ala Leu Arg Gln Ala Ala Met Arg His Glu Glu Asn Glu Arg
245 250 255

Ala Phe Phe Glu Arg Leu Gly Thr Ile Ser Asp Arg Tyr Glu Ala Ser
260 265 270

Pro Asp Phe Thr Glu Tyr Ser Ala Val Ser Asn Ile Gly Tyr Gly Phe
275 280 285

Ile Gln Gln His Asp Phe Pro Gly Leu Ala Ile Asn Asp Asn Leu Gln
290 295 300

Asp Ala Asn Gln Ile Gln Leu Tyr His Gly Ala Pro Tyr Ile Phe Thr
305 310 315 320

Phe Gly Asp Val Asp Lys His Asn Gln Gln
325 330

<210> 33

<211> 235

<212> PRT

<213> Citrobacter rodentium

<400> 33

SB-0008.ST25.txt

Met Arg Pro Thr Ser Leu Asn Leu Thr Leu Pro Ser Leu Pro Leu Pro
1 5 10 15

Ser Ser Ser Asn Ser Ile Ser Ala Thr Asp Ile Gln Ser Leu Val Lys
20 25 30

Met Ser Gly Val Arg Trp Val Lys Asn Asn Gln Gln Leu Cys Phe His
35 40 45

Gly Thr Asp Leu Lys Ile Tyr Gln His Leu Glu Ala Ala Leu Asp Lys
50 55 60

Ile Glu Ser Thr Asp Thr Gly Arg Thr Leu Leu Asn Cys Ile Glu Leu
65 70 75 80

Thr Ser Arg Leu Lys Ser Glu Lys Leu Ala Ile His Leu Asp Ser Ala
85 90 95

Glu Leu Gly Val Ile Ala His Cys Asn Ala Asp Ala Glu Asn Ser Arg
100 105 110

Gly Thr Gly Ser Asp Phe His Cys Asn Leu Asn Ala Val Glu Tyr Pro
115 120 125

Cys Gly Gln Gly Ile Ser Leu Val Asp Phe His Ala Cys Ile Val Phe
130 135 140

His Glu Leu Leu His Val Phe His Asn Leu Asn Gly Glu Arg Leu Lys
145 150 155 160

Val Glu Ser Ser Gln Pro Glu Leu Gln Thr His Ser Pro Leu Leu Leu
165 170 175

Glu Glu Ala Arg Thr Val Gly Leu Gly Ala Phe Ser Glu Glu Val Leu
180 185 190

Ser Glu Asn Lys Phe Arg Glu Glu Ile Gly Met Pro Arg Arg Thr Phe
195 200 205

Tyr Pro His Asp Ser Ser Leu Ile His Asp Asp Asn Thr Val Thr Gln
210 215 220

Arg Phe Gln Arg Lys Lys Leu His Pro Leu Leu
225 230 235

<210> 34

<211> 232

<212> PRT

<213> Enteropathogenic E. coli

<400> 34

SB-0008.ST25.txt

Met Arg Pro Thr Ser Leu Asn Leu Val Leu His Gln Ser Ser Thr Ser
 1 5 10 15

Ser Ser Met Ser Asp Thr Asp Ile Glu Ser Leu Val Lys Ala Ser Ser
 20 25 30

Val Gln Trp Ile Lys Asn Asn Pro Gln Leu Arg Phe Gln Gly Thr Asp
 35 40 45

His Asn Ile Tyr Gln Gln Ile Glu Ala Ala Leu Asp Lys Ile Gly Ser
 50 55 60

Thr Glu Thr Gly Arg Val Leu Leu Asn Ala Ile Glu Ser Ile Ser Arg
 65 70 75 80

Leu Lys Ser Glu Thr Val Val Ile His Leu Asn Ser Ser Arg Leu Gly
 85 90 95

Val Met Ala His Arg Asp Ile Asp Ala Glu Asn His Arg Gly Thr Gly
 100 105 110

Ser Asp Phe His Cys Asn Leu Asn Ala Val Glu Tyr Pro Cys Gly Glu
 115 120 125

Gly Ile Ser Val Val Asp Phe His Ala Thr Ile Val Phe His Glu Leu
 130 135 140

Leu His Val Phe His Asn Leu Asn Gly Glu Arg Leu Lys Val Glu Ser
 145 150 155 160

Ser Arg Pro Glu Ser Gln Lys Tyr Ser Pro Leu Leu Leu Glu Glu Ala
 165 170 175

Arg Thr Val Gly Leu Gly Ala Phe Ser Glu Glu Val Leu Ser Glu Asn
 180 185 190

Lys Phe Arg Glu Glu Ile Gly Met Pro Arg Arg Thr Ser Tyr Pro His
 195 200 205

Asp Ser Ala Leu Ile His Asp Asp Asn Thr Val Ser Leu Gly Phe Gln
 210 215 220

Gln Val Arg Leu His Pro Leu Leu
 225 230

<210> 35
 <211> 232
 <212> PRT
 <213> Enterohemorrhagic E. coli

<220>
 <221> MISC_FEATURE

SB-0008.ST25.txt

<222> (208)..(208)

<223> Xaa = Arg or His

<400> 35

Met Arg Pro Thr Ser Leu Asn Leu Val Leu His Gln Ser Ser Arg Ser
 1 5 10 15

Ser Ser Met Ser Asp Thr Asp Ile Glu Ser Leu Val Lys Ala Ser Ser
 20 25 30

Val Gln Trp Ile Lys Asn Asn Pro Gln Leu Arg Phe Gln Gly Thr Asp
 35 40 45

His Asn Ile Tyr Gln Gln Ile Glu Ala Ala Leu Asp Lys Ile Gly Ser
 50 55 60

Thr Glu Thr Gly Arg Val Leu Leu Asn Ala Ile Glu Ser Ile Ser Arg
 65 70 75 80

Leu Lys Ser Glu Thr Val Val Ile His Leu Asn Ser Ser Arg Leu Gly
 85 90 95

Val Met Ala His Arg Asp Ile Asp Ala Glu Asn His Arg Gly Thr Gly
 100 105 110

Ser Asp Phe His Cys Asn Leu Asn Ala Val Glu Tyr Pro Cys Gly Glu
 115 120 125

Gly Ile Ser Val Val Asp Phe His Ala Thr Ile Val Phe His Glu Leu
 130 135 140

Leu His Val Phe His Asn Leu Asn Gly Glu Arg Leu Lys Val Glu Ser
 145 150 155 160

Ser Arg Ala Glu Ser Gln Lys Tyr Ser Pro Leu Leu Leu Glu Glu Ala
 165 170 175

Arg Thr Val Gly Leu Gly Ala Phe Ser Glu Glu Val Leu Ser Glu Asn
 180 185 190

Lys Phe His Glu Glu Ile Gly Met Pro Arg Arg Thr Ser Tyr Pro Xaa
 195 200 205

Asp Ser Ala Leu Ile His Asp Asp Asn Thr Val Ser Leu Gly Phe Gln
 210 215 220

Gln Val Arg Leu His Pro Leu Leu
 225 230

<210> 36

<211> 168

<212> PRT

SB-0008.ST25.txt

<213> Citrobacter rodentium

<400> 36

Tyr Phe Asn Glu Ser Pro Asn Val Tyr Asp Lys Lys Tyr Ile Ser Gly
 1 5 10 15

Val Thr Arg Gly Val Ala Glu Leu Lys Gln Glu Gly Phe Ile Asn Glu
 20 25 30

Lys Ala Arg Arg Leu Ala Tyr Met Gln Ala Met Tyr Ser Val Cys Pro
 35 40 45

Glu Glu Phe Lys Pro Ile Ser Arg Asn Glu Ala Ser Thr Pro Glu Gly
 50 55 60

Ser Trp Leu Thr Val Ile Ser Gly Lys Arg Pro Met Gly Gln Phe Ser
 65 70 75 80

Val Asp Ser Leu Tyr His Pro Asp Leu His Ala Leu Cys Glu Leu Pro
 85 90 95

Asp Ile Cys Cys Lys Ile Phe Pro Lys Glu Asn Asn Asp Phe Leu Tyr
 100 105 110

Ile Val Ile Val Tyr Arg Asn Asp Ser Pro Leu Gly Glu Gln Arg Ala
 115 120 125

Asn Arg Phe Ile Glu Leu Tyr Asn Ile Lys Arg Asp Ile Met Gln Glu
 130 135 140

Leu Asn Tyr Glu Ser Pro Glu Leu Lys Ala Val Lys Ser Glu Met Ile
 145 150 155 160

Ile Ala Arg Glu Met Gly Glu Ile
 165

<210> 37

<211> 154

<212> PRT

<213> Citrobacter rodentium

<400> 37

Asn Val Tyr Asp Lys Lys Tyr Ile Ser Gly Val Thr Arg Gly Val Ala
 1 5 10 15

Glu Leu Lys Gln Glu Gly Phe Ile Asn Glu Lys Ala Arg Arg Leu Ala
 20 25 30

Tyr Met Gln Ala Met Tyr Ser Val Cys Pro Glu Glu Phe Lys Pro Ile
 35 40 45

Ser Arg Asn Glu Ala Ser Thr Pro Glu Gly Ser Trp Leu Thr Val Ile
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50

55

60

Ser Gly Lys Arg Pro Met Gly Gln Phe Ser Val Asp Ser Leu Tyr His
 65 70 75 80

Pro Asp Leu His Ala Leu Cys Glu Leu Pro Asp Ile Cys Cys Lys Ile
 85 90 95

Phe Pro Lys Glu Asn Asn Asp Phe Leu Tyr Ile Val Ile Val Tyr Arg
 100 105 110

Asn Asp Ser Pro Leu Gly Glu Gln Arg Ala Asn Arg Phe Ile Glu Leu
 115 120 125

Tyr Asn Ile Lys Arg Asp Ile Met Gln Glu Leu Asn Tyr Glu Ser Pro
 130 135 140

Glu Leu Lys Ala Val Lys Ser Glu Met Ile
 145 150

<210> 38

<211> 224

<212> PRT

<213> Enteropathogenic E. coli

<400> 38

Met Ile Asn Pro Val Thr Asn Thr Gln Gly Val Ser Pro Ile Asn Thr
 1 5 10 15

Lys Tyr Ala Glu His Val Val Lys Asn Ile Tyr Pro Lys Ile Lys His
 20 25 30

Asp Tyr Phe Asn Glu Ser Pro Asn Ile Tyr Asp Lys Lys Tyr Ile Ser
 35 40 45

Gly Ile Thr Arg Gly Val Ala Glu Leu Lys Gln Glu Glu Phe Val Asn
 50 55 60

Glu Lys Ala Arg Arg Phe Ser Tyr Met Lys Thr Met Tyr Ser Val Cys
 65 70 75 80

Pro Glu Ala Phe Glu Pro Ile Ser Arg Asn Glu Ala Ser Thr Pro Glu
 85 90 95

Gly Ser Trp Leu Thr Val Ile Ser Gly Lys Arg Pro Met Gly Gln Phe
 100 105 110

Ser Val Asp Ser Leu Tyr Asn Pro Asp Leu His Ala Leu Cys Glu Leu
 115 120 125

Pro Asp Ile Cys Cys Lys Ile Phe Pro Lys Glu Asn Asn Asp Phe Leu
 130 135 140

SB-0008.ST25.txt

Tyr Ile Val Val Val Tyr Arg Asn Asp Ser Pro Leu Gly Glu Gln Arg
145 150 155 160

Ala Asn Arg Phe Ile Glu Leu Tyr Asn Ile Lys Arg Asp Ile Met Gln
165 170 175

Glu Leu Asn Tyr Glu Leu Pro Glu Leu Lys Ala Val Lys Ser Glu Met
180 185 190

Ile Ile Ala Arg Glu Met Gly Glu Ile Phe Ser Tyr Met Pro Gly Glu
195 200 205

Ile Asp Ser Tyr Met Lys Tyr Ile Asn Asn Lys Leu Ser Lys Ile Glu
210 215 220

<210> 39

<211> 224

<212> PRT

<213> Enterohemorrhagic E. coli

<400> 39

Met Ile Asn Pro Val Thr Asn Thr Gln Gly Val Ser Pro Ile Asn Thr
1 5 10 15

Lys Tyr Ala Glu His Val Val Lys Asn Ile Tyr Pro Glu Ile Lys His
20 25 30

Asp Tyr Phe Asn Glu Ser Pro Asn Ile Tyr Asp Lys Lys Tyr Ile Ser
35 40 45

Gly Ile Thr Arg Gly Val Ala Glu Leu Lys Gln Glu Glu Phe Val Asn
50 55 60

Glu Lys Ala Arg Arg Phe Ser Tyr Met Lys Thr Met Tyr Ser Val Cys
65 70 75 80

Pro Glu Ala Phe Glu Pro Ile Ser Arg Asn Glu Ala Ser Thr Pro Glu
85 90 95

Gly Ser Trp Leu Thr Val Ile Ser Gly Lys Arg Pro Met Gly Gln Phe
100 105 110

Ser Val Asp Ser Leu Tyr Asn Pro Asp Leu His Ala Leu Cys Glu Leu
115 120 125

Pro Asp Ile Cys Cys Lys Ile Phe Pro Lys Glu Asn Asn Asp Phe Leu
130 135 140

Tyr Ile Val Val Val Tyr Arg Asn Asp Ser Pro Leu Gly Glu Gln Arg
145 150 155 160

SB-0008.ST25.txt

Ala Asn Arg Phe Ile Glu Leu Tyr Asn Ile Lys Arg Asp Ile Met Gln
165 170 175

Glu Leu Asn Tyr Glu Leu Pro Glu Leu Lys Ala Val Lys Ser Glu Met
180 185 190

Ile Ile Ala Arg Glu Met Gly Glu Ile Phe Ser Tyr Met Pro Gly Glu
195 200 205

Ile Asp Ser Tyr Met Lys Tyr Ile Asn Asn Lys Leu Ser Lys Ile Glu
210 215 220

<210> 40

<211> 188

<212> PRT

<213> Citrobacter rodentium

<400> 40

Met Leu Pro Thr Ser Gly Ser Ser Ala Asn Leu Tyr Ser Trp Met Tyr
1 5 10 15

Ile Ser Gly Lys Glu Asn Pro Ser Thr Pro Glu Ser Val Ser Glu Leu
20 25 30

Asn His Asn His Phe Leu Ser Pro Glu Leu Gln Glu Lys Leu Asp Val
35 40 45

Met Phe Ala Ile Tyr Ser Cys Ala Arg Asn Asn Asp Glu Arg Glu Asn
50 55 60

Ile Tyr Pro Glu Leu Arg Asp Phe Val Ser Ser Leu Met Asp Lys Arg
65 70 75 80

Asn Asn Val Phe Glu Val Ile Asn Glu Asp Thr Asp Glu Val Thr Gly
85 90 95

Ala Leu Arg Ala Gly Met Thr Ile Glu Asp Arg Asp Ser Tyr Ile Arg
100 105 110

Asp Leu Phe Phe Leu His Ser Leu Lys Val Lys Ile Glu Glu Ser Arg
115 120 125

Gln Asp Lys Glu Asp Trp Lys Cys Lys Val Tyr Asp Leu Leu Cys Pro
130 135 140

His His Ser Ser Glu Leu Tyr Gly Asp Leu Arg Ala Ile Lys Cys Leu
145 150 155 160

Val Glu Gly Cys Ser Asp Asp Phe Ser Pro Phe Asp Thr Ile Lys Val
165 170 175

SB-0008.ST25.txt

Pro Asp Leu Thr Tyr Asn Lys Gly Ser Leu Gln Cys
 180 185

<210> 41
 <211> 171
 <212> PRT
 <213> Citrobacter rodentium

<400> 41

Ala Asn Leu Tyr Ser Trp Met Tyr Ile Ser Gly Lys Glu Asn Pro Ser
 1 5 10 15

Thr Pro Glu Ser Val Ser Glu Leu Asn His Asn His Phe Leu Ser Pro
 20 25 30

Glu Leu Gln Glu Lys Leu Asp Val Met Phe Ala Ile Tyr Ser Cys Ala
 35 40 45

Arg Asn Asn Asp Glu Arg Glu Asn Ile Tyr Pro Glu Leu Arg Asp Phe
 50 55 60

Val Ser Ser Leu Met Asp Lys Arg Asn Asn Val Phe Glu Val Ile Asn
 65 70 75 80

Glu Asp Thr Asp Glu Val Thr Gly Ala Leu Arg Ala Gly Met Thr Ile
 85 90 95

Glu Asp Arg Asp Ser Tyr Ile Arg Asp Leu Phe Phe Leu His Ser Leu
 100 105 110

Lys Val Lys Ile Glu Glu Ser Arg Gln Asp Lys Glu Asp Trp Lys Cys
 115 120 125

Lys Val Tyr Asp Leu Leu Cys Pro His His Ser Ser Glu Leu Tyr Gly
 130 135 140

Asp Leu Arg Ala Ile Lys Cys Leu Val Glu Gly Cys Ser Asp Asp Phe
 145 150 155 160

Ser Pro Phe Asp Thr Ile Lys Val Pro Asp Leu
 165 170

<210> 42
 <211> 189
 <212> PRT
 <213> Enteropathogenic E. coli

<400> 42

Met Leu Pro Thr Ser Gly Ser Ser Ala Asn Leu Tyr Ser Trp Met Tyr
 1 5 10 15

Val Ser Gly Arg Gly Asn Pro Ser Thr Pro Glu Ser Val Ser Glu Leu
 20 25 30

SB-0008.ST25.txt

Asn His Asn His Phe Leu Ser Pro Glu Leu Gln Asp Lys Leu Asp Val
35 40 45

Met Val Ser Ile Tyr Ser Cys Ala Arg Asn Asn Glu Leu Glu Glu
50 55 60

Ile Phe Gln Glu Leu Ser Ala Phe Val Ser Gly Leu Met Asp Lys Arg
65 70 75 80

Asn Ser Val Phe Glu Val Arg Asn Glu Asn Thr Asp Glu Val Val Gly
85 90 95

Ala Leu Arg Ala Gly Met Thr Ile Glu Asp Arg Asp Ser Tyr Ile Arg
100 105 110

Asp Leu Phe Phe Leu His Ser Leu Lys Val Lys Ile Glu Glu Ser Arg
115 120 125

Gln Gly Lys Glu Asp Ser Lys Cys Lys Val Tyr Asn Leu Leu Cys Pro
130 135 140

His His Ser Ser Glu Leu Tyr Gly Asp Leu Arg Ala Met Lys Cys Leu
145 150 155 160

Val Glu Gly Cys Ser Asp Asp Phe Asn Pro Phe Asp Ile Ile Arg Val
165 170 175

Pro Asp Leu Thr Tyr Asn Lys Gly Ser Leu Gln Cys Gly
180 185

<210> 43

<211> 189

<212> PRT

<213> Enterohemorrhagic E. coli

<400> 43

Met Leu Pro Thr Ser Gly Ser Ser Ala Asn Leu Tyr Ser Trp Met Tyr
1 5 10 15

Val Ser Gly Arg Gly Asn Pro Ser Thr Pro Glu Ser Val Ser Glu Leu
20 25 30

Asn His Asn His Phe Leu Ser Pro Glu Leu Gln Asp Lys Leu Asp Val
35 40 45

Met Val Ser Ile Tyr Ser Cys Ala Arg Asn Asn Glu Leu Glu Glu
50 55 60

Ile Phe Gln Glu Leu Ser Ala Phe Val Ser Gly Leu Met Asp Lys Arg
65 70 75 80

SB-0008.ST25.txt

Asn Ser Val Phe Glu Val Arg Asn Glu Asn Thr Asp Glu Val Val Gly
85 90 95

Ala Leu Arg Ala Gly Met Thr Ile Glu Asp Arg Asp Ser Tyr Ile Arg
100 105 110

Asp Leu Phe Phe Leu His Ser Leu Lys Val Lys Ile Glu Glu Ser Arg
115 120 125

Gln Gly Lys Glu Asp Ser Lys Cys Lys Val Tyr Asn Leu Leu Cys Pro
130 135 140

His His Ser Ser Glu Leu Tyr Gly Asp Leu Arg Ala Met Lys Cys Leu
145 150 155 160

Val Glu Gly Cys Ser Asp Asp Phe Asn Pro Phe Asp Ile Ile Arg Val
165 170 175

Pro Asp Leu Thr Tyr Asn Lys Gly Ser Leu Gln Cys Gly
180 185

<210> 44
<211> 40
<212> DNA
<213> Artificial Sequence

<220>
<223> primer Z6024F

<400> 44
agatctgaag gagatattat gaacattcaa ccgaccatac 40

<210> 45
<211> 34
<212> DNA
<213> Artificial Sequence

<220>
<223> primer Z6024R

<400> 45
ctcgaggact cttgtttctt cgattatatac aaag 34

<210> 46
<211> 28
<212> DNA
<213> Artificial Sequence

<220>
<223> primer NT10

<400> 46
ccggtaacctc taaccattga cgcactcg 28

<210> 47
<211> 29
<212> DNA

SB-0008.ST25.txt

<213> Artificial Sequence

<220>

<223> primer NT11

<400> 47

aacctgcaga actaggtatc tctaatgcc

29

<210> 48

<211> 29

<212> DNA

<213> Artificial Sequence

<220>

<223> primer NT12

<400> 48

aacctgcagc tgactatcct cgtatatgg

29

<210> 49

<211> 27

<212> DNA

<213> Artificial Sequence

<220>

<223> primer NT13

<400> 49

ccgagcttag gtaatgagac tgtcagc

27

<210> 50

<211> 22

<212> DNA

<213> Artificial Sequence

<220>

<223> primer del1F

<400> 50

ggtaccacca cacagaataa tc

22

<210> 51

<211> 26

<212> DNA

<213> Artificial Sequence

<220>

<223> primer del1R

<400> 51

cgcttagccta tatactgctg ttggtt

26

<210> 52

<211> 28

<212> DNA

<213> Artificial Sequence

<220>

<223> primer del12F

<400> 52

gctagctgac aggcaactct tggactgg

28

SB-0008.ST25.txt

<210> 53
<211> 29
<212> DNA
<213> Artificial Sequence

<220>
<223> primer del2R

<400> 53
gagctcaaca taatggatg gattatgtat

29

<210> 54
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<223> primer

<400> 54
ttccatatatga acattcaacc gacc

24

<210> 55
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<223> primer

<400> 55
ggaattcaat aatagctgcc atcc

24

<210> 56
<211> 135
<212> PRT
<213> Salmonella

<400> 56

Met Glu Ser Lys Asn Ser Asp Tyr Val Ile Pro Asp Ser Val Lys Asn
1 5 10 15

Tyr Asn Gly Glu Pro Leu Tyr Ile Leu Val Ser Leu Trp Cys Lys Leu
20 25 30

Gln Glu Lys Trp Ile Ser Arg Asn Asp Ile Ala Glu Ala Phe Gly Ile
35 40 45

Asn Leu Arg Arg Ala Ser Phe Ile Ile Thr Tyr Ile Ser Arg Arg Lys
50 55 60

Glu Lys Ile Ser Phe Arg Val Arg Tyr Val Ser Tyr Gly Asn Leu His
65 70 75 80

Tyr Lys Arg Leu Glu Ile Phe Ile Tyr Asn Val Asn Leu Glu Ala Ala
85 90 95

SB-0008.ST25.txt

Pro Thr Glu Ser His Val Ser Thr Gly Pro Lys Arg Lys Thr Leu Arg
 100 105 110

Val Gly Asn Gly Ile Val Gly Gln Ser Ser Ile Trp Asn Glu Met Ile
 115 120 125

Met Arg Arg Lys Lys Glu Ser
 130 135

<210> 57
 <211> 131
 <212> PRT
 <213> Enterobacteriaceae

<400> 57

Met Cys Glu Gly Tyr Val Glu Lys Pro Leu Tyr Leu Leu Ile Ala Glu
 1 5 10 15

Trp Met Met Ala Glu Asn Arg Trp Val Ile Ala Arg Glu Ile Ser Ile
 20 25 30

His Phe Asp Ile Glu His Ser Lys Ala Val Asn Thr Leu Thr Tyr Ile
 35 40 45

Leu Ser Glu Val Thr Glu Ile Ser Cys Glu Val Lys Met Ile Pro Asn
 50 55 60

Lys Leu Glu Gly Arg Gly Cys Gln Cys Gln Arg Leu Val Lys Val Val
 65 70 75 80

Asp Ile Asp Glu Gln Ile Tyr Ala Arg Leu Arg Asn Asn Ser Arg Glu
 85 90 95

Lys Leu Val Gly Val Arg Lys Thr Pro Arg Ile Pro Ala Val Pro Leu
 100 105 110

Thr Glu Leu Asn Arg Glu Gln Lys Trp Gln Met Met Leu Ser Lys Ser
 115 120 125

Met Arg Arg
 130

<210> 58
 <211> 170
 <212> PRT
 <213> Citrobacter rodentium

<400> 58

Met Cys Pro Asp Asn Thr His Ala Lys Lys Gln Tyr Leu Thr Pro Gly
 1 5 10 15

Asn Asp Ile His Tyr Pro Gly Gln Thr Asn His Asp Ala Cys Phe Ile
 20 25 30

SB-0008.ST25.txt

Pro Val Ser Val Arg Gln Tyr Ala Gly Glu Pro Leu Tyr Ile Ile Val
35 40 45

Ala His Trp Cys Leu Leu Gln Gln Asn Trp Val Gln Arg Asn Gln Ile
50 55 60

Ala Glu Ala Phe His Ile Thr Ala Arg Arg Ala Ser Tyr Leu Ile Ala
65 70 75 80

Tyr Leu Arg Ser Lys Thr Ser Arg Val Val Ser Ile Cys Arg His Gln
85 90 95

Thr Leu Pro Asn Lys Ala Arg Arg Tyr Glu Ile Tyr Val Ile Arg Val
100 105 110

Leu Asp Ser Pro Thr Pro Ser Thr Arg Arg Glu Lys Ala Gly Pro Pro
115 120 125

Leu Val Ser Lys Arg Arg Val Gly Asn Gly Asp Arg Ser Met Ala Asn
130 135 140

Glu Leu Trp Asn Arg Leu Cys Ser Asn Arg Asn Ala Gly Lys Ile Leu
145 150 155 160

Lys Lys Lys Glu Asp Glu Asp Asp Gly Thr
165 170

<210> 59
<211> 12
<212> PRT
<213> Citrobacter rodentium

<220>
<221> MISC_FEATURE
<222> (9)..(9)
<223> Xaa = Ile or Leu

<400> 59

Gln Gln Glu Asn Ala Pro Ser Ser Xaa Gln Thr Arg
1 5 10

<210> 60
<211> 981
<212> DNA
<213> Enterohemorrhagic E. coli

<400> 60
atgcttcac cgataaggac aactttccat aactcagtaa atatagtgc gagttcaccc 60
tgtcaaacgg tttctttgc agggaaaggaa tatgagttaa aggtcattga tgaaaaaacg 120
cctattctt ttcatgtgtt tgaacctaat cctgaacgtat ataagaaaga tgaggttcca 180
atagttata ctaagcagca tccctattta gataatgtca caaatgcggc aaggatagag 240

SB-0008.ST25.txt

agtgatcgta tgataggtat tttgttgat ggcgatttt cagtcaacca aaagactgct	300
ttttcaaaat tggAACgaga tttgaaaat gtaatgataa tctatcggga agatgttgac	360
ttcagtatgt atgacagaaa actatcatg atttatcatg atattatatg tgaacaaagg	420
ttacgaactg aagacaaaag agatgaatac ttgttgaatc tggtagagaa agagctgagg	480
gaaatttcaa aggCGCAGGA ttcttgatt tctatgtatg caaagaaaag aaatcatgca	540
tggtttGatt tcttcagaaa tttagccta ttaaaAGCAG gagagatatt caggtgcaca	600
tataatacaa agaatcacgg tatttcattc ggggagggggt gtatctatct tgatatggat	660
atgatactta caggtaagct tggTACAATA tatgctcctg atggaatttc aatgcATGtG	720
gatcgTCgtA atgatagtgt aaatattgaa aatagtGCAA taattgttaa ccgtagtaat	780
catccTgCTC tacttgaggg actttttttt atgcataGTA aagttagatgc tcAtCCatAT	840
tatgatggtt tggggAAAGG agttaagaaa tattttAATT ttacaccatt acataattat	900
aatcatTTTt gtgactttat tgagttAAC caccctaata taatcatgaa cacaagtcag	960
tatacatgca gttcatggta a	981

<210> 61
<211> 531
<212> DNA
<213> Enterohemorrhagic E. coli

<400> 61 atgaatgtcc ttcgagctca agtagcatct agcggtcgag gggagttac attaggtaat	60
gagactgtca gcattgtatt taatgaaacc gatggcgTT ttctatccag cggcagttagt	120
gggggattgc ttactgagtt attcctttat gggttaata acggccctga agctcttcgc	180
gataggatgc tcagtatgct ttccgactca ggtgaagcac aatcgcaaga gagtattcag	240
gacaaaatat ctcaatgtaa gttccctgtt agttcaggaa atttccagtg cccgccagag	300
tctattcagt gtccaattac actagagaga cccgaagaag gagtgtttgt caaaaattca	360
gatagttcgg cagtatgctg cttatttGAT tttgatgcat tttctcgTTT agctagtGAA	420
ggctcatatc atccactgac ccgagaacca ataacggcat caatgattat aagtccTgAT	480
aaatgtgttt atgatcctat caagggAAAC ttcattataa aagatagtta a	531

<210> 62
<211> 912
<212> DNA
<213> Enterohemorrhagic E. coli

<400> 62 atgttatcgc cctttctat aaatttggga tggatcgta attcttaac cagaaacctg	60
acttcgcctg ataatcgtgt tttatcctct gtaaggatg ctgctgttca ctctgatagc	120
gggacgcaag taacggTTGG caacagaaca tatcgTTTG tggTCACTGA taataagttt	180
tgcgttacaa gagaaagtca tagtggttgt tttactaATC tggTGCACAG gttgggatgg	240
cctaaggag agattagcag aaaaattgag gctatgctga atacatcgcc agtgagcACG	300

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actatagaaa	gaggcttgt	tcattcgAAC	agacCTGATT	tacCCAGT	ggATTATGCG	360
cagccggagt	tacCTCCAGC	ggATTATACT	caatCAGAGT	tgCCGAGGGT	tagCAACAA	420
aaatCACCCG	tgCCAGGTAA	tgttATTGgt	aaAGGTGGTA	atGCTGTCgt	gtATGAAGAT	480
atGGAAGATA	caACAAAAGT	gttGAAGATG	tttACTATAT	ctCAAAGCCA	tGAAGAGGTG	540
acaAGCAG	ttCGTTGTTT	caATCAGTAT	tatGGTCCG	ggAGTGcaga	gAAAATATA	600
aatGATAATG	gaaATGTTAT	tggTATTAGA	atGAATAAAA	taaATGGGGA	atCTTTTG	660
gatATTCCAT	cattACCAGC	acaAGCTGAA	caggCTATT	acGATATGTT	tgACAGACTG	720
gagaAAAAAG	gaATTCTTT	tgttgataca	acAGAAACAA	atGTTTATA	tgATCGTATG	780
agAAATGAAT	ttaATCCAAT	agATATATCA	tcttATAATG	tttCTGATAT	ttCATGGAGT	840
gaACATCAAG	tcatGCAATC	ttatCACGGA	ggAAAGCTGG	atCTTATTAG	tgtAGTATTA	900
agTAAGATAT	aa					912

<210> 63
<211> 882
<212> DNA
<213> Enterohemorrhagic E. coli

<400> 63	atgttatcgc	catattctgt	aaatttggga	tgttcatgga	attcttaac	cagaaacctg	60
	acttcgcctg	ataatcgtgt	tttacctct	gtaagggatg	ctgccgttca	ttctgataat	120
	ggggcgcaag	taaaggttgg	caacagaaca	tatcgtgtt	ttgccaccga	taataagttt	180
	tgcgttacaa	gagaaagtca	tagtgggtgt	tttactaattc	tgttgcacag	gctggatgg	240
	cctaaggggg	agattagcag	gaaaattgag	gtcatgctga	atgcattcacc	agtgagcgct	300
	gctatggaaa	gaggcattgt	tcattcgAAC	agacCTGATT	tacCCAGT	tgattatgca	360
	ccgcccagagt	taccgagggt	ggactataac	aggTTGTCAG	tacCTGGTAA	tgttattggc	420
	aaagggggga	acgctgttagt	atatgaagat	gctgaggatg	caACAAAAGT	cctgaagatg	480
	tttactacat	ctcaaagcaa	tgaagagggt	acaAGCAG	ttcgttgctt	caACCAATAT	540
	tatggtccg	ggagtgcaga	aaaaatata	ggcaataatg	gtgatattat	tggTATTAGA	600
	atggataaaa	taaATGGAGA	atcgctttt	aatattcgt	ccttgccagc	acaggctgag	660
	catgctattt	acgatATGTT	tgatagactg	gagCAAAAAG	gaATTCTTT	tgtcgataca	720
	acagagacaa	atgtcttata	tgaccgcgcg	aagaatgagt	ttaATCCAAT	agatATATCA	780
	tcttATAATG	tttCCGACCG	ttCATGGAGT	gaaAGTCaaa	taatGCAATC	ttatCATGGC	840
	ggaaagcaag	atCTTATTAG	tgtggtatta	agtaaaaattt	ag		882

<210> 64
<211> 153
<212> DNA
<213> Enterohemorrhagic E. coli

<400> 64	atggtaatgc	ctggattagt	atcatatata	tcatcgactt	cattcgCGAA	tgagatggcg	60
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gagatgcgtc agcaggtaat ggaagggcag attgggtggat ttctcctggg aggggagaga	120
gttagagttt cttatttatt tcaattgcat taa	153

<210> 65
<211> 576
<212> DNA
<213> Enterohemorrhagic E. coli

<400> 65 atgccattaa cctcagatata tagatcacat tcatttaatc ttgggggtgga gggtgttcgt	60
gccccgaattt tagccaatgg gcgcggagat attacagtgc gtggtaaaac tgctcgttatt	120
gtgtatgatt ctactaatgg gcgcgtttca tccagtgccg gtaatggcg attgcgttct	180
gagttattgc ttttgggatt taatagtggt cctcgagccc ttggtgagag aatgctaagt	240
atgcgttcgg actcaggtga agcacaatcg caagagagta ttcagaacaa aatatctcaa	300
tgttaagttt ctgtttgtcc agagagactt cagtgcgcgc ttggaggctat tcagtgtcca	360
attacactgg agcagcctga aaaaggtatt tttgtgaaga attcagatgg ttcagatgtta	420
tgtactttat ttgatgccgc tgcattttct cgtttgggtt gtgaaggctt accccaccca	480
ctgaccggg aaccaataac ggcataata attgtaaaac atgaagaatg catttatgac	540
gataccagag gaaaccttcat tataaagggt aattga	576

<210> 66
<211> 630
<212> DNA
<213> Enterohemorrhagic E. coli

<220>
<221> misc_feature
<222> (439)..(439)
<223> n = any nucleotide

<400> 66 atgcctgtta ccacctaag tatcccaagt atatctcaat tatctcctgc aagagtacag	60
tctttcagg atgcagccag acttggaaat ggaataagaa tatccattgg tagtggccaa	120
tattctgttc actatgtcca actactggat ggattttcag ttgaaccgggt gagaggaggc	180
ttactggata ggctattggg gcgtgagcat cgaatggata gaagggtgt ggctctggaa	240
aggcaattaa atggaggtgt cgattttta agtagtgtta ataactattt tcagagtgtc	300
atggcagaac acagagaaaa taaaacaggt aataaaatata taatggaaaa aataaattct	360
tgtgtatgg gaacggattc taatcacttt tcttgcccg agtcatttt gacatgcccg	420
ataacgctgg acacaccta gactggagtg ttcatgagaa actcacgagg tgctgagata	480
tgctctctat atgataagga tgcgttagtgc caacttgttgc aaactgggttgc aactcatcct	540
ctgagtcgag aacctataac agaatcaatg attatgagaa aagacgaatg tcactttgtat	600
gcaaaaaagag aagcttttg ttgttaagtga	630

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<210> 67
 <211> 642
 <212> DNA
 <213> Enterohemorrhagic E. coli

<400> 67
 atgcctgttag attaacgcc ttatattta cctgggtta gtttttgc tgacattcct 60
 caagaaacct tgtctgagat acgtaatcag actattcgtg gagaagctca agtaagactg 120
 ggtgagttga tggtgtcaat acgacctatg caggtaaatg gatattttat ggaaagtctt 180
 aaccaggatg gtttatcgaa tgataacatc cagattggcc ttcaatatat agaacatatt 240
 gaacgtacac ttaatcatgg tagttgaca agccgtgaag ttacagtact gcgtgaaatt 300
 gagatgctcg aaaatatgga attgcttct aactaccagt tagaggagtt gttagataaa 360
 attgaagtat gtgcatttaa tggagcat gcacaattgc aagtgccaga gagcttacga 420
 acatgccctg ttacattatg tgaaccagaa gatgggtat ttatgaggaa ttcaatgaat 480
 tcaaatgttt gtatgttga tgataaaatg tcattaatat atcttgttaa aacaaggcg 540
 gctcatcctt tgagcagggta atcaatcgca gttcaatga ttgttaggaag agataattgt 600
 gctttgact ctgacagagg taacttcgtt ttaaaaaatt aa 642

<210> 68
 <211> 642
 <212> DNA
 <213> Enterohemorrhagic E. coli

<400> 68
 atgcctgttag attaacgcc ttatattta cctgggtta gtttttgc tgacattcct 60
 caagaaacct tgtctgagat acgtaatcag actattcgtg gagaagctca aataagactg 120
 ggtgagttga tggtgtcaat acgacctatg caggtaaatg gatattttat ggaaagtctt 180
 aaccaggatg gtttatcgaa tgataatatc cagattggcc ttcaatatat agaacatatt 240
 gaacgtacac ttaatcatgg tagttgaca agccgtgaag ttacagtact gcgtgaaatt 300
 gagatgctcg aaaatatgga ttgcttct aactaccagt tagaggagtt gttagataaa 360
 attgaagtat gtgcatttaa tggagcat gcacaattgc aagtgccaga gagcttacga 420
 acatgccctg ttacattatg tgaaccagaa gatgggtat ttatgaggaa ttcaatgaat 480
 tcaaatgttt gtatgttga tgataaaatg gcattaatac atcttgttaa aacaaggcg 540
 gctcatcctt tgagcagggta atcaatcgca gttcaatga ttgttaggaag agataattgt 600
 gctttgacc ctgacagagg taacttcgtt ttaaaaaatt aa 642

<210> 69
 <211> 630
 <212> DNA
 <213> Enterohemorrhagic E. coli

<400> 69
 atgcctgtta ccacctaag tatcccaagt atatctcaat tatctcctgc aggagtacag 60
 tctttgcagg atgctgccag actgaaagt ggaataagaa tatccattgg tagtgccaa 120

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tattctgttc	actatgtcca	gctactggat	ggattttcag	ttgaaccgggt	gagaggaggc	180
ttactggata	ggctattggg	gcgtgagcat	cgaatggaga	gaagggctgt	ggctctggaa	240
aggcaattaa	atggaggtgt	cgattttta	agttagtta	ataactattt	tcagagtgtc	300
atggcagaac	acagagaaaa	taaaaacaagt	aataaaatat	taatggaaaa	aataaattct	360
tgtttattta	gacctgattc	taatcacttt	tcttgcccg	agtcatttt	gacatccccg	420
ataacgctgg	acacaccta	gactggggtg	ttcatgagaa	actcacgagg	tgctgagata	480
tgctcttat	atgataagga	cgcgttagt	caacttgg	aaactgg	agctcatcct	540
ctgagtcgag	aacctataac	agaatcaatg	attatgagaa	aagatgaatg	tcactttgat	600
acaaaaagag	aagcttttg	ttgtaagtga				630
<210>	70					
<211>	576					
<212>	DNA					
<213>	Enterohemorrhagic E. coli					
<400>	70					
atgccattaa	cctcagat	at tagatcacat	tcatttaatc	ttgggggtgga	ggttgttcgt	60
gcccgaattt	tagccaatgg	gcgcggagat	attacagt	gtggtaaaac	tgtcagtatt	120
gtgtatgatt	ctactaatgg	gcgcctttca	tccagtg	gtaatggcg	attgtttct	180
gagttattgc	ttttgggatt	taatagtgg	cctcgagccc	ttggtgagag	aatgtaagt	240
atgcttcgg	actcaggtga	agcacaatcg	caagagagta	ttcagaacaa	aatatctcaa	300
tgttaagttt	ctgtttgtcc	agagagactt	cagtccccgc	ttgaggctat	tcartgtcca	360
attacactgg	agcagcctga	aaaaggatt	tttgtaaga	attcagatgg	ttcagatgt	420
tgtactttat	ttgatgccgc	tgcattttct	cgtttgg	gtgaaggctt	accccaccca	480
ctgaccggg	aaccaataac	ggcatcaata	attgtaaaac	atgaagaatg	cattatgac	540
gataccagag	gaaacttcgt	tataaagggt	aattga			576
<210>	71					
<211>	510					
<212>	DNA					
<213>	Enterohemorrhagic E. coli					
<400>	71					
atggacgctt	ttattgtaga	tcctgttcaa	gggaaactat	attcgggtt	aagccataca	60
gaactagccg	atatcattag	attggctgat	tctgtgaaa	atcaattgaa	tggaggcaat	120
tcatttctt	atgtattcag	tacatatatg	ggcaggtt	tttctgaatt	tatgcata	180
aatgataaca	gaattgaatt	tttacagcgg	cgattacatt	catgttcatt	tttagttaat	240
attgaagaaa	tgtcttacat	agatgaagca	ttacagt	cgattacgct	ggcaattcct	300
caacgaggtg	tttttttaag	aatgctgaa	ggttccagag	tatgttagttt	atatgatgaa	360
atggctctt	ctcgataat	taatgatgg	atgcatacacc	cactaagcag	agagccaata	420
acattatcaa	tgcttgtggc	cagagagcag	tgtgagttt	attgcagtat	cggtcacttt	480

acggtgagga gtgatttta ttcagtgtag

510

<210> 72
<211> 231
<212> DNA
<213> Enterohemorrhagic E. coli

<400> 72
atggcagacc gcaaacagca ccgcgtatc gcggagcgtc gtcacatcca gactgaaatc 60
aaccgcagac tttcccgcbc atcacgcgtc gcgcaaatca tgcacatcaa tatgctgcat 120
gagcgcagcc acgcactatc aaacatttat tccgcctctg ttttcagcta tctggcggat 180
gatctgcacg agtttcaaca gctcatccag cagcaaaaca aactccattta a 231

<210> 73
<211> 176
<212> PRT
<213> Enterohemorrhagic E. coli

<400> 73

Met	Asn	Val	Leu	Arg	Ala	Gln	Val	Ala	Ser	Ser	Gly	Arg	Gly	Glu	Phe
1														10	15

Thr	Leu	Gly	Asn	Glu	Thr	Val	Ser	Ile	Val	Phe	Asn	Glu	Thr	Asp	Gly
														20	25

Arg	Phe	Leu	Ser	Ser	Gly	Ser	Ser	Gly	Gly	Leu	Leu	Thr	Glu	Leu	Phe
														35	40

Leu	Tyr	Gly	Phe	Asn	Asn	Gly	Pro	Glu	Ala	Leu	Arg	Asp	Arg	Met	Leu
														50	55

Ser	Met	Leu	Ser	Asp	Ser	Gly	Glu	Ala	Gln	Ser	Gln	Glu	Ser	Ile	Gln
														65	70

Asp	Lys	Ile	Ser	Gln	Cys	Lys	Phe	Pro	Val	Ser	Ser	Gly	Asn	Phe	Gln
														85	90

Cys	Pro	Pro	Glu	Ser	Ile	Gln	Cys	Pro	Ile	Thr	Leu	Glu	Arg	Pro	Glu
														100	105

Glu	Gly	Val	Phe	Val	Lys	Asn	Ser	Asp	Ser	Ser	Ala	Val	Cys	Cys	Leu
														115	120

Phe	Asp	Phe	Asp	Ala	Phe	Ser	Arg	Leu	Ala	Ser	Glu	Gly	Ser	Tyr	His
														130	135

Pro	Leu	Thr	Arg	Glu	Pro	Ile	Thr	Ala	Ser	Met	Ile	Ile	Ser	Pro	Asp
														145	150

Lys	Cys	Val	Tyr	Asp	Pro	Ile	Lys	Gly	Asn	Phe	Ile	Ile	Lys	Asp	Ser
														165	170

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<210> 74
<211> 303
<212> PRT
<213> Enterohemorrhagic E. coli

<400> 74

Met Leu Ser Pro Ser Ser Ile Asn Leu Gly Cys Ser Trp Asn Ser Leu
1 5 10 15

Thr Arg Asn Leu Thr Ser Pro Asp Asn Arg Val Leu Ser Ser Val Arg
20 25 30

Asp Ala Ala Val His Ser Asp Ser Gly Thr Gln Val Thr Val Gly Asn
35 40 45

Arg Thr Tyr Arg Val Val Val Thr Asp Asn Lys Phe Cys Val Thr Arg
50 55 60

Glu Ser His Ser Gly Cys Phe Thr Asn Leu Leu His Arg Leu Gly Trp
65 70 75 80

Pro Lys Gly Glu Ile Ser Arg Lys Ile Glu Ala Met Leu Asn Thr Ser
85 90 95

Pro Val Ser Thr Thr Ile Glu Arg Gly Ser Val His Ser Asn Arg Pro
100 105 110

Asp Leu Pro Pro Val Asp Tyr Ala Gln Pro Glu Leu Pro Pro Ala Asp
115 120 125

Tyr Thr Gln Ser Glu Leu Pro Arg Val Ser Asn Asn Lys Ser Pro Val
130 135 140

Pro Gly Asn Val Ile Gly Lys Gly Asn Ala Val Val Tyr Glu Asp
145 150 155 160

Met Glu Asp Thr Thr Lys Val Leu Lys Met Phe Thr Ile Ser Gln Ser
165 170 175

His Glu Glu Val Thr Ser Glu Val Arg Cys Phe Asn Gln Tyr Tyr Gly
180 185 190

Ser Gly Ser Ala Glu Lys Ile Tyr Asn Asp Asn Gly Asn Val Ile Gly
195 200 205

Ile Arg Met Asn Lys Ile Asn Gly Glu Ser Leu Leu Asp Ile Pro Ser
210 215 220

Leu Pro Ala Gln Ala Glu Gln Ala Ile Tyr Asp Met Phe Asp Arg Leu
225 230 235 240

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Glu Lys Lys Gly Ile Leu Phe Val Asp Thr Thr Glu Thr Asn Val Leu
 245 250 255

Tyr Asp Arg Met Arg Asn Glu Phe Asn Pro Ile Asp Ile Ser Ser Tyr
 260 265 270

Asn Val Ser Asp Ile Ser Trp Ser Glu His Gln Val Met Gln Ser Tyr
 275 280 285

His Gly Gly Lys Leu Asp Leu Ile Ser Val Val Leu Ser Lys Ile
 290 295 300

<210> 75

<211> 293

<212> PRT

<213> Enterohemorrhagic E. coli

<400> 75

Met Leu Ser Pro Tyr Ser Val Asn Leu Gly Cys Ser Trp Asn Ser Leu
 1 5 10 15

Thr Arg Asn Leu Thr Ser Pro Asp Asn Arg Val Leu Ser Ser Val Arg
 20 25 30

Asp Ala Ala Val His Ser Asp Asn Gly Ala Gln Val Lys Val Gly Asn
 35 40 45

Arg Thr Tyr Arg Val Val Ala Thr Asp Asn Lys Phe Cys Val Thr Arg
 50 55 60

Glu Ser His Ser Gly Cys Phe Thr Asn Leu Leu His Arg Leu Gly Trp
 65 70 75 80

Pro Lys Gly Glu Ile Ser Arg Lys Ile Glu Val Met Leu Asn Ala Ser
 85 90 95

Pro Val Ser Ala Ala Met Glu Arg Gly Ile Val His Ser Asn Arg Pro
 100 105 110

Asp Leu Pro Pro Val Asp Tyr Ala Pro Pro Glu Leu Pro Ser Val Asp
 115 120 125

Tyr Asn Arg Leu Ser Val Pro Gly Asn Val Ile Gly Lys Gly Gly Asn
 130 135 140

Ala Val Val Tyr Glu Asp Ala Glu Asp Ala Thr Lys Val Leu Lys Met
 145 150 155 160

Phe Thr Thr Ser Gln Ser Asn Glu Glu Val Thr Ser Glu Val Arg Cys
 165 170 175

Phe Asn Gln Tyr Tyr Gly Ala Gly Ser Ala Glu Lys Ile Tyr Gly Asn
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185

190

Asn Gly Asp Ile Ile Gly Ile Arg Met Asp Lys Ile Asn Gly Glu Ser
195 200 205

Leu Leu Asn Ile Ser Ser Leu Pro Ala Gln Ala Glu His Ala Ile Tyr
210 215 220

Asp Met Phe Asp Arg Leu Glu Gln Lys Gly Ile Leu Phe Val Asp Thr
225 230 235 240

Thr Glu Thr Asn Val Leu Tyr Asp Arg Ala Lys Asn Glu Phe Asn Pro
245 250 255

Ile Asp Ile Ser Ser Tyr Asn Val Ser Asp Arg Ser Trp Ser Glu Ser
260 265 270

Gln Ile Met Gln Ser Tyr His Gly Gly Lys Gln Asp Leu Ile Ser val
275 280 285

val Leu Ser Lys Ile
290

<210> 76

<211> 50

<212> PRT

<213> Enterohemorrhagic E. coli

<400> 76

Met Val Met Pro Gly Leu Val Ser Tyr Ile Ser Ser Thr Ser Phe Ala
1 5 10 15

Asn Glu Met Ala Glu Met Arg Gln Gln Val Met Glu Gly Gln Ile Gly
20 25 30

Gly Phe Leu Leu Gly Gly Glu Arg Val Arg Val Ser Tyr Leu Phe Gln
35 40 45

Leu His
50

<210> 77

<211> 191

<212> PRT

<213> Enterohemorrhagic E. coli

<400> 77

Met Pro Leu Thr Ser Asp Ile Arg Ser His Ser Phe Asn Leu Gly Val
1 5 10 15

Glu Val Val Arg Ala Arg Ile Val Ala Asn Gly Arg Gly Asp Ile Thr
20 25 30

SB-0008.ST25.txt

Val Gly Gly Glu Thr Val Ser Ile Val Tyr Asp Ser Thr Asn Gly Arg
35 40 45

Phe Ser Ser Ser Gly Gly Asn Gly Leu Leu Ser Glu Leu Leu Leu
50 55 60

Leu Gly Phe Asn Ser Gly Pro Arg Ala Leu Gly Glu Arg Met Leu Ser
65 70 75 80

Met Leu Ser Asp Ser Gly Glu Ala Gln Ser Gln Glu Ser Ile Gln Asn
85 90 95

Lys Ile Ser Gln Cys Lys Phe Ser Val Cys Pro Glu Arg Leu Gln Cys
100 105 110

Pro Leu Glu Ala Ile Gln Cys Pro Ile Thr Leu Glu Gln Pro Glu Lys
115 120 125

Gly Ile Phe Val Lys Asn Ser Asp Gly Ser Asp Val Cys Thr Leu Phe
130 135 140

Asp Ala Ala Ala Phe Ser Arg Leu Val Gly Glu Gly Leu Pro His Pro
145 150 155 160

Leu Thr Arg Glu Pro Ile Thr Ala Ser Ile Ile Val Lys His Glu Glu
165 170 175

Cys Ile Tyr Asp Asp Thr Arg Gly Asn Phe Ile Ile Lys Gly Asn
180 185 190

<210> 78

<211> 209

<212> PRT

<213> Enterohemorrhagic E. coli

<220>

<221> MISC_FEATURE

<222> (147)..(147)

<223> Xaa = any amino acid

<400> 78

Met Pro Val Thr Thr Leu Ser Ile Pro Ser Ile Ser Gln Leu Ser Pro
1 5 10 15

Ala Arg Val Gln Ser Leu Gln Asp Ala Ala Arg Leu Glu Ser Gly Ile
20 25 30

Arg Ile Ser Ile Gly Ser Gly Gln Tyr Ser Val His Tyr Val Gln Leu
35 40 45

Leu Asp Gly Phe Ser Val Glu Pro Val Arg Gly Gly Leu Leu Asp Arg
50 55 60

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Leu Leu Gly Arg Glu His Arg Met Asp Arg Arg Ala Val Ala Leu Glu
65 70 75 80

Arg Gln Leu Asn Gly Gly Val Asp Phe Leu Ser Ser Val Asn Asn Tyr
85 90 95

Phe Gln Ser Val Met Ala Glu His Arg Glu Asn Lys Thr Gly Asn Lys
100 105 110

Ile Leu Met Glu Lys Ile Asn Ser Cys Val Phe Gly Thr Asp Ser Asn
115 120 125

His Phe Ser Cys Pro Glu Ser Phe Leu Thr Cys Pro Ile Thr Leu Asp
130 135 140

Thr Pro Xaa Thr Gly Val Phe Met Arg Asn Ser Arg Gly Ala Glu Ile
145 150 155 160

Cys Ser Leu Tyr Asp Lys Asp Ala Leu Val Gln Leu Val Glu Thr Gly
165 170 175

Gly Thr His Pro Leu Ser Arg Glu Pro Ile Thr Glu Ser Met Ile Met
180 185 190

Arg Lys Asp Glu Cys His Phe Asp Ala Lys Arg Glu Ala Phe Cys Cys
195 200 205

Lys

<210> 79

<211> 213

<212> PRT

<213> Enterohemorrhagic E. coli

<400> 79

Met Pro Val Asp Leu Thr Pro Tyr Ile Leu Pro Gly Val Ser Phe Leu
1 5 10 15

Ser Asp Ile Pro Gln Glu Thr Leu Ser Glu Ile Arg Asn Gln Thr Ile
20 25 30

Arg Gly Glu Ala Gln Val Arg Leu Gly Glu Leu Met Val Ser Ile Arg
35 40 45

Pro Met Gln Val Asn Gly Tyr Phe Met Gly Ser Leu Asn Gln Asp Gly
50 55 60

Leu Ser Asn Asp Asn Ile Gln Ile Gly Leu Gln Tyr Ile Glu His Ile
65 70 75 80

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Glu Arg Thr Leu Asn His Gly Ser Leu Thr Ser Arg Glu Val Thr Val
85 90 95

Leu Arg Glu Ile Glu Met Leu Glu Asn Met Glu Leu Leu Ser Asn Tyr
100 105 110

Gln Leu Glu Glu Leu Leu Asp Lys Ile Glu Val Cys Ala Phe Asn Val
115 120 125

Glu His Ala Gln Leu Gln Val Pro Glu Ser Leu Arg Thr Cys Pro Val
130 135 140

Thr Leu Cys Glu Pro Glu Asp Gly Val Phe Met Arg Asn Ser Met Asn
145 150 155 160

Ser Asn Val Cys Met Leu Tyr Asp Lys Met Ser Leu Ile Tyr Leu Val
165 170 175

Lys Thr Arg Ala Ala His Pro Leu Ser Arg Glu Ser Ile Ala Val Ser
180 185 190

Met Ile Val Gly Arg Asp Asn Cys Ala Phe Asp Ser Asp Arg Gly Asn
195 200 205

Phe Val Leu Lys Asn
210

<210> 80

<211> 213

<212> PRT

<213> Enterohemorrhagic E. coli

<400> 80

Met Pro Val Asp Leu Thr Pro Tyr Ile Leu Pro Gly Val Ser Phe Leu
1 5 10 15

Ser Asp Ile Pro Gln Glu Thr Leu Ser Glu Ile Arg Asn Gln Thr Ile
20 25 30

Arg Gly Glu Ala Gln Ile Arg Leu Gly Glu Leu Met Val Ser Ile Arg
35 40 45

Pro Met Gln Val Asn Gly Tyr Phe Met Gly Ser Leu Asn Gln Asp Gly
50 55 60

Leu Ser Asn Asp Asn Ile Gln Ile Gly Leu Gln Tyr Ile Glu His Ile
65 70 75 80

Glu Arg Thr Leu Asn His Gly Ser Leu Thr Ser Arg Glu Val Thr Val
85 90 95

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Leu Arg Glu Ile Glu Met Leu Glu Asn Met Asp Leu Leu Ser Asn Tyr
100 105 110

Gln Leu Glu Glu Leu Leu Asp Lys Ile Glu Val Cys Ala Phe Asn Val
115 120 125

Glu His Ala Gln Leu Gln Val Pro Glu Ser Leu Arg Thr Cys Pro Val
130 135 140 145

Thr Leu Cys Glu Pro Glu Asp Gly Val Phe Met Arg Asn Ser Met Asn
145 150 155 160

Ser Asn Val Cys Met Leu Tyr Asp Lys Met Ala Leu Ile His Leu Val
165 170 175

Lys Thr Arg Ala Ala His Pro Leu Ser Arg Glu Ser Ile Ala Val Ser
180 185 190

Met Ile Val Gly Arg Asp Asn Cys Ala Phe Asp Pro Asp Arg Gly Asn
195 200 205

Phe Val Leu Lys Asn
210

<210> 81
<211> 209
<212> PRT
<213> Enterohemorrhagic E. coli

<400> 81

Met Pro Val Thr Thr Leu Ser Ile Pro Ser Ile Ser Gln Leu Ser Pro
1 5 10 15

Ala Gly Val Gln Ser Leu Gln Asp Ala Ala Arg Leu Glu Ser Gly Ile
20 25 30

Arg Ile Ser Ile Gly Ser Gly Gln Tyr Ser Val His Tyr Val Gln Leu
35 40 45

Leu Asp Gly Phe Ser Val Glu Pro Val Arg Gly Gly Leu Leu Asp Arg
50 55 60

Leu Leu Gly Arg Glu His Arg Met Glu Arg Arg Ala Val Ala Leu Glu
65 70 75 80

Arg Gln Leu Asn Gly Gly Val Asp Phe Leu Ser Ser Val Asn Asn Tyr
85 90 95

Phe Gln Ser Val Met Ala Glu His Arg Glu Asn Lys Thr Ser Asn Lys
100 105 110

Ile Leu Met Glu Lys Ile Asn Ser Cys Leu Phe Arg Pro Asp Ser Asn
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115

120

125

His Phe Ser Cys Pro Glu Ser Phe Leu Thr Cys Pro Ile Thr Leu Asp
 130 135 140

Thr Pro Glu Thr Gly Val Phe Met Arg Asn Ser Arg Gly Ala Glu Ile
 145 150 155 160

Cys Ser Leu Tyr Asp Lys Asp Ala Leu Val Gln Leu Val Glu Thr Gly
 165 170 175

Gly Ala His Pro Leu Ser Arg Glu Pro Ile Thr Glu Ser Met Ile Met
 180 185 190

Arg Lys Asp Glu Cys His Phe Asp Thr Lys Arg Glu Ala Phe Cys Cys
 195 200 205

Lys

<210> 82

<211> 191

<212> PRT

<213> Enterohemorrhagic E. coli

<400> 82

Met Pro Leu Thr Ser Asp Ile Arg Ser His Ser Phe Asn Leu Gly Val
 1 5 10 15

Glu Val Val Arg Ala Arg Ile Val Ala Asn Gly Arg Gly Asp Ile Thr
 20 25 30

Val Gly Gly Glu Thr Val Ser Ile Val Tyr Asp Ser Thr Asn Gly Arg
 35 40 45

Phe Ser Ser Ser Gly Gly Asn Gly Leu Leu Ser Glu Leu Leu Leu
 50 55 60

Leu Gly Phe Asn Ser Gly Pro Arg Ala Leu Gly Glu Arg Met Leu Ser
 65 70 75 80

Met Leu Ser Asp Ser Gly Glu Ala Gln Ser Gln Glu Ser Ile Gln Asn
 85 90 95

Lys Ile Ser Gln Cys Lys Phe Ser Val Cys Pro Glu Arg Leu Gln Cys
 100 105 110

Pro Leu Glu Ala Ile Gln Cys Pro Ile Thr Leu Glu Gln Pro Glu Lys
 115 120 125

Gly Ile Phe Val Lys Asn Ser Asp Gly Ser Asp Val Cys Thr Leu Phe
 130 135 140

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Asp Ala Ala Ala Phe Ser Arg Leu Val Gly Glu Gly Leu Pro His Pro
145 150 155 160

Leu Thr Arg Glu Pro Ile Thr Ala Ser Ile Ile Val Lys His Glu Glu
165 170 175

Cys Ile Tyr Asp Asp Thr Arg Gly Asn Phe Val Ile Lys Gly Asn
180 185 190

<210> 83

<211> 169

<212> PRT

<213> Enterohemorrhagic E. coli

<400> 83

Met Asp Ala Phe Ile Val Asp Pro Val Gln Gly Glu Leu Tyr Ser Gly
1 5 10 15

Leu Ser His Thr Glu Leu Ala Asp Ile Ile Arg Leu Ala Asp Ser Val
20 25 30

Glu Asn Gln Leu Asn Gly Gly Asn Ser Phe Leu Asp Val Phe Ser Thr
35 40 45

Tyr Met Gly Gln Val Ile Ser Glu Phe Met His Ser Asn Asp Asn Arg
50 55 60

Ile Glu Leu Leu Gln Arg Arg Leu His Ser Cys Ser Phe Leu Val Asn
65 70 75 80

Ile Glu Glu Met Ser Tyr Ile Asp Glu Ala Leu Gln Cys Pro Ile Thr
85 90 95

Leu Ala Ile Pro Gln Arg Gly Val Phe Leu Arg Asn Ala Glu Gly Ser
100 105 110

Arg Val Cys Ser Leu Tyr Asp Glu Met Ala Leu Ser Arg Ile Ile Asn
115 120 125

Asp Gly Met His His Pro Leu Ser Arg Glu Pro Ile Thr Leu Ser Met
130 135 140

Leu Val Ala Arg Glu Gln Cys Glu Phe Asp Cys Ser Ile Gly His Phe
145 150 155 160

Thr Val Arg Ser Asp Cys Tyr Ser Val
165

<210> 84

<211> 76

<212> PRT

SB-0008.ST25.txt

<213> Enterohemorrhagic E. coli

<400> 84

Met Ala Asp Arg Lys Gln His Arg Ala Ile Ala Glu Arg Arg His Ile
1 5 10 15

Gln Thr Glu Ile Asn Arg Arg Leu Ser Arg Ala Ser Arg Val Ala Gln
20 25 30

Ile Met His Ile Asn Met Leu His Glu Arg Ser His Ala Leu Ser Asn
35 40 45

Ile Tyr Ser Ala Ser Val Phe Ser Tyr Leu Ala Asp Asp Leu His Glu
50 55 60

Phe Gln Gln Leu Ile Gln Gln Asn Lys Leu His
65 70 75